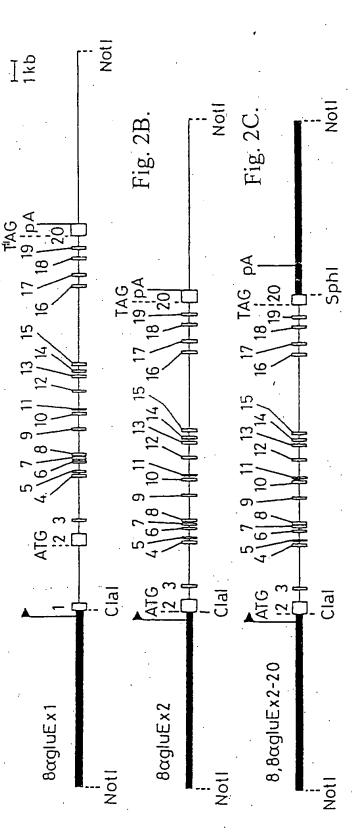




Fig. 2.A



Transcription Initiation site.

 $lpha_{\mathrm{S1}}$  casein sequence, promoler or  $3^{l}$  untranslated region.

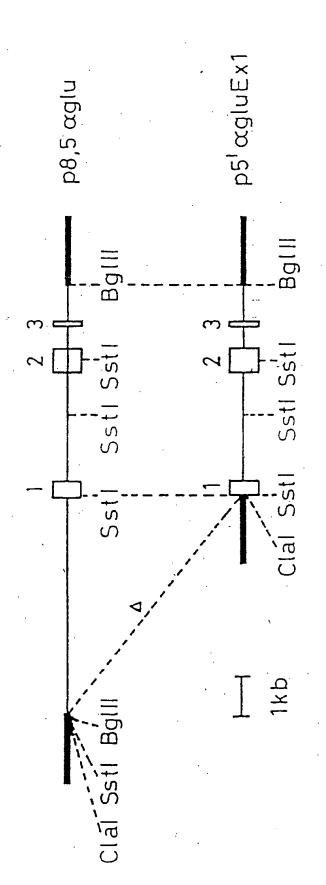
The boxes represent the exons in the  $\alpha\text{--}glucosidase$  sequence, the thin line represents he intron sequences.

The numbers above the boxes are the exon numbers

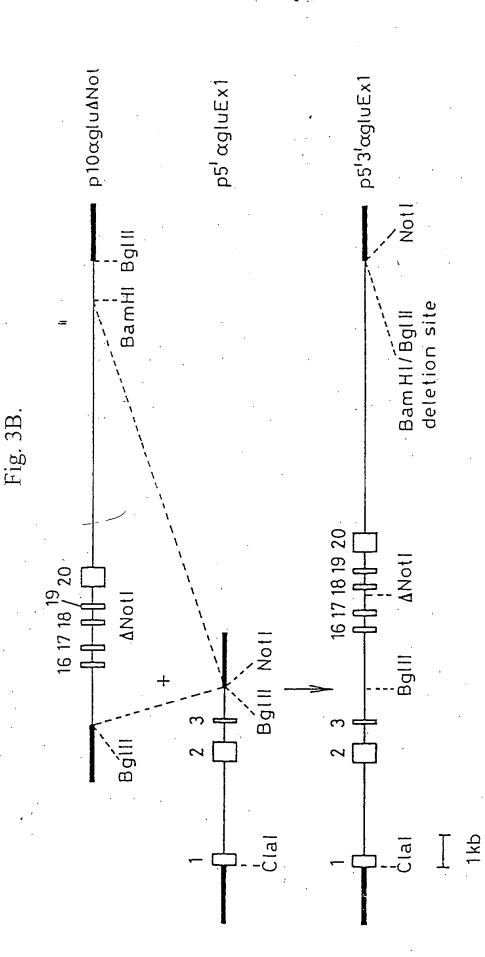
= polyadeny lation signal. βĄ

ATG = translation initiation site. TAG = translation stop codon

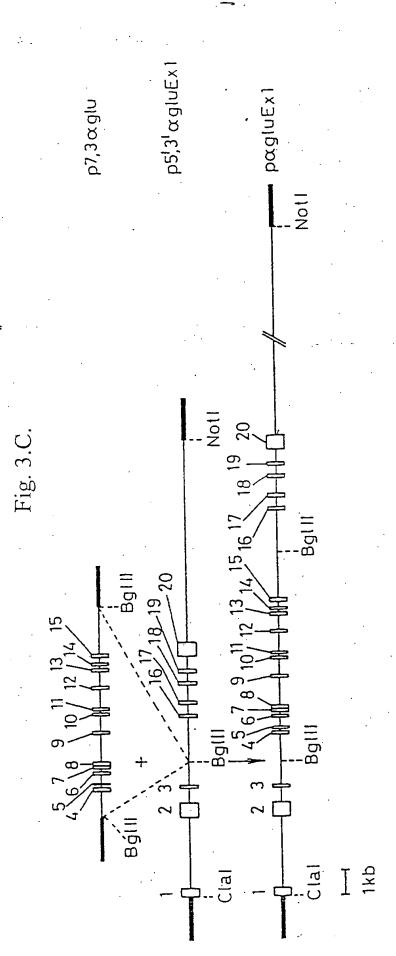
Fig. 3A.



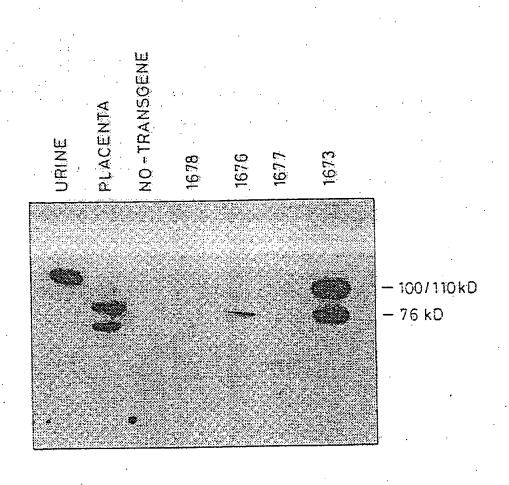
= pKUN vector sequence = intron  $\alpha$ -glu = exon  $\alpha$ -glu

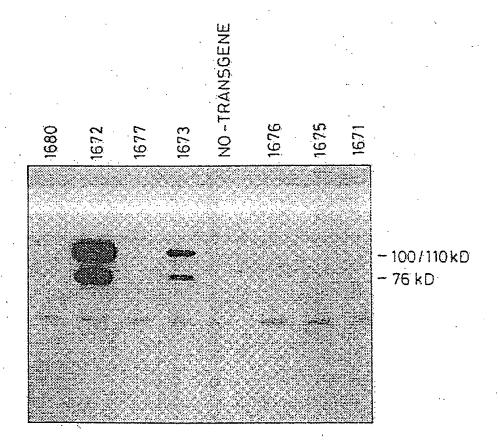


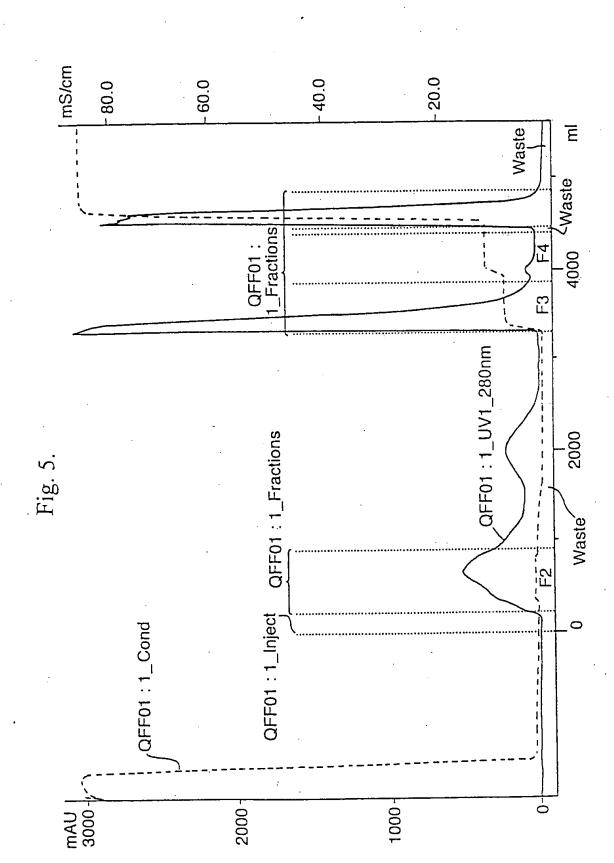
= pKUN vector sequence -= intron  $\alpha$ -glu □ = exon α-glu

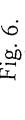


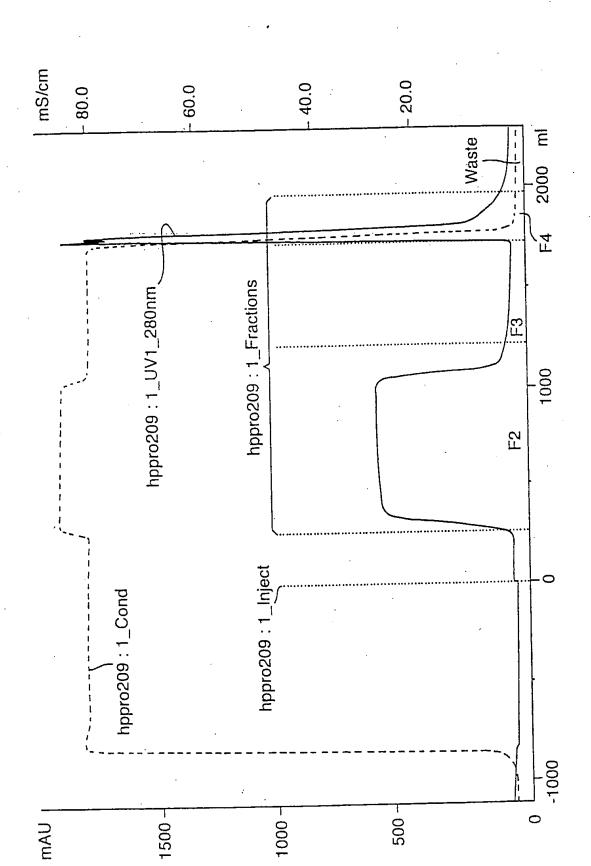
= pKUN vector sequence -= intron α-glu 🛚 = exon α-glu











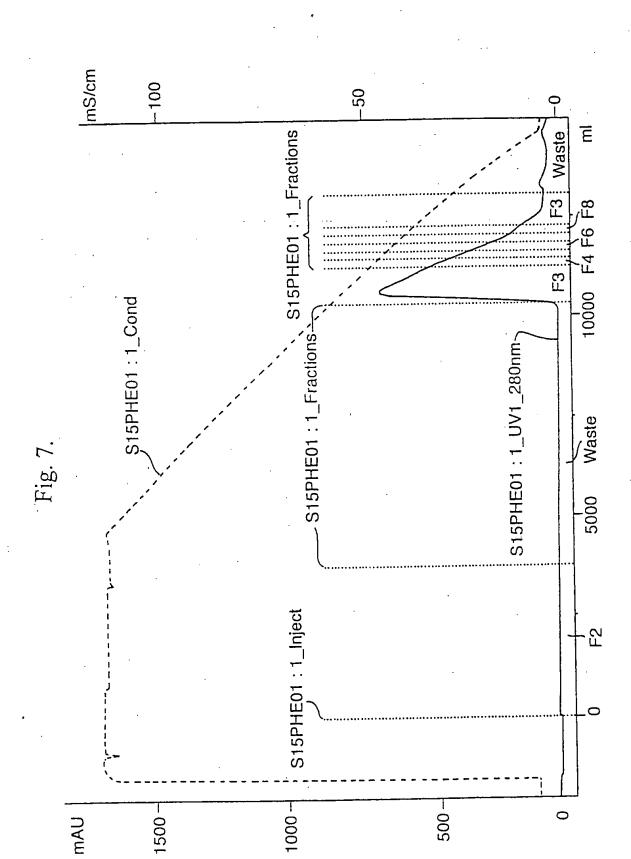


Fig. 8.

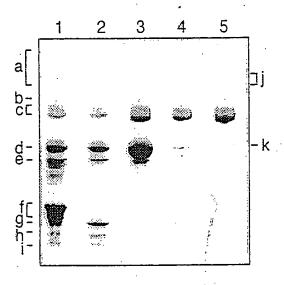


Fig. 9.

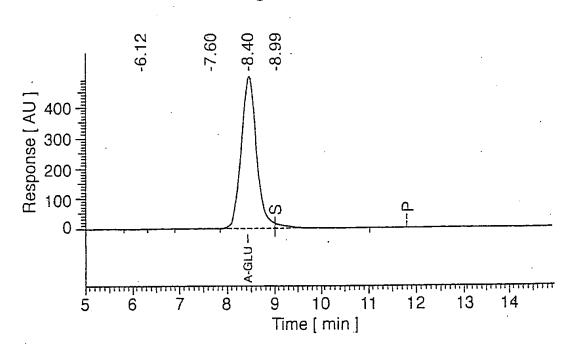


Fig. 10.

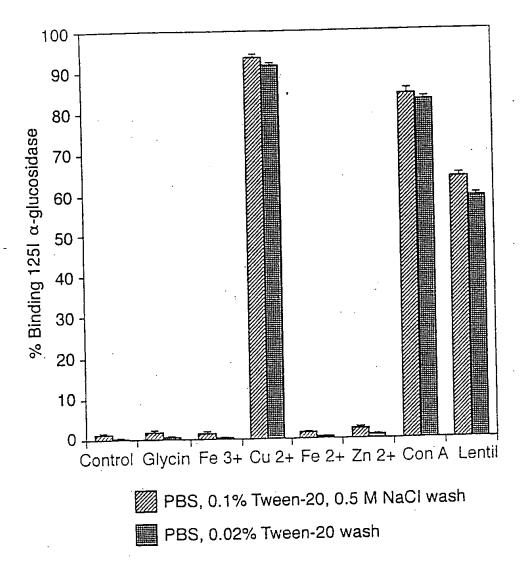


Fig. 11. A.

AU 1M AmSO 110 kD 76 kD 76 kD 0.0000 0 M AmSO 0 M AmSO 110 kD 0 M AmSO 110 kD 11

Fig. 11. B.

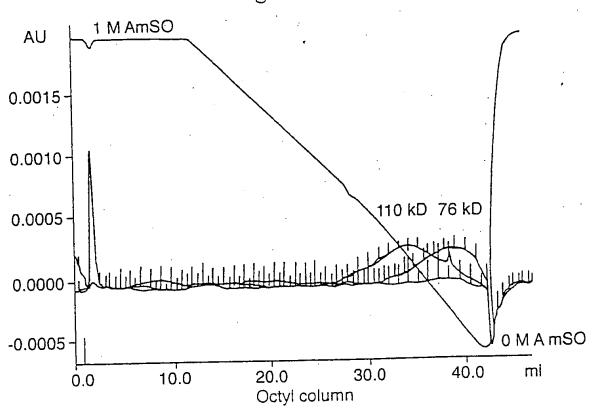
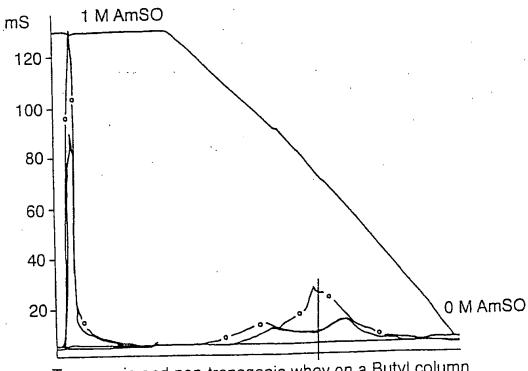
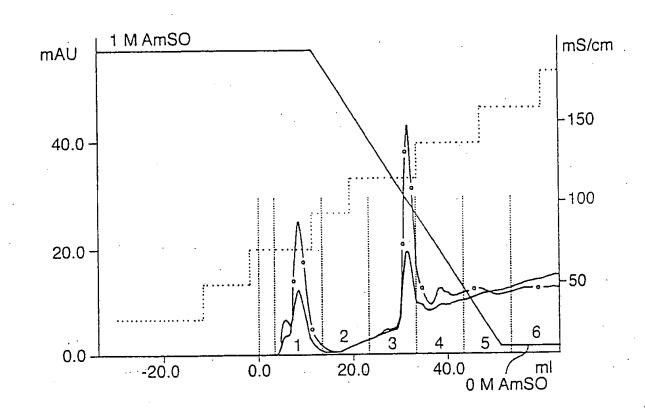


Fig. 11. C.



Transgenic and non-transgenic whey on a Butyl column

Fig. 11. D.



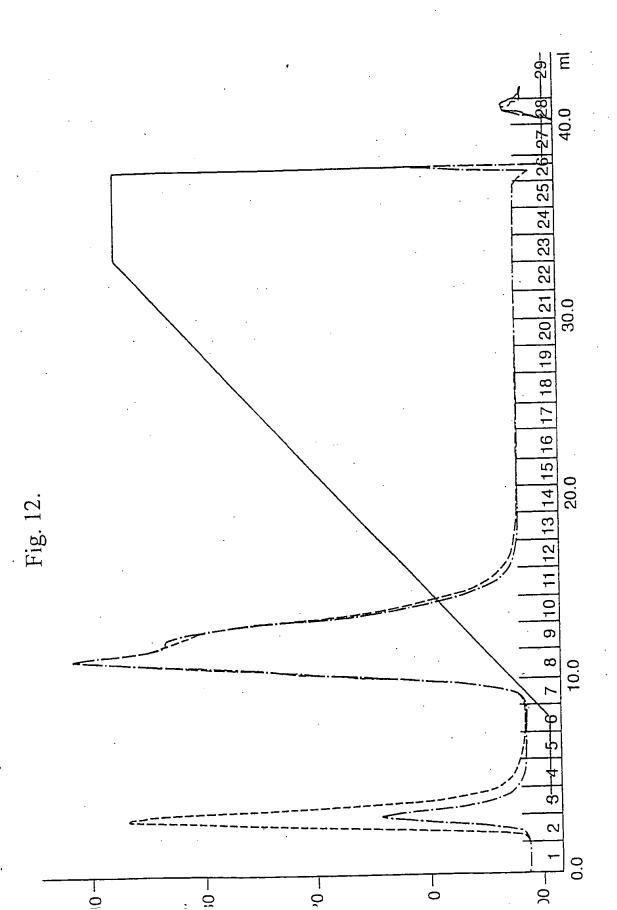


Fig. 13. A.

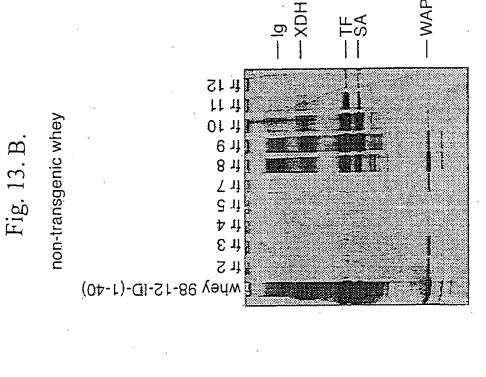
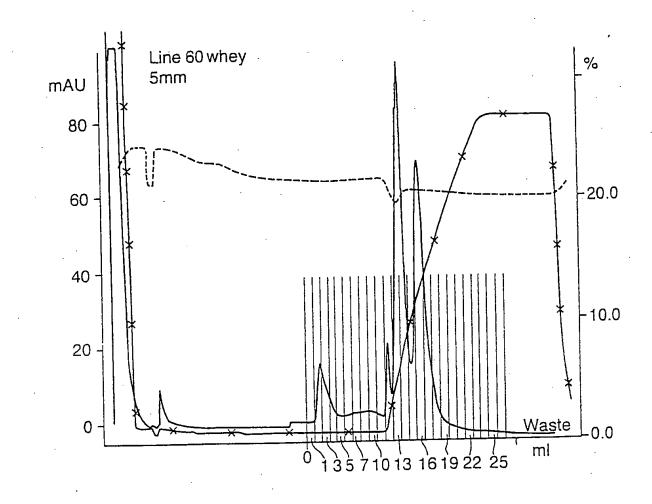


Fig. 14.



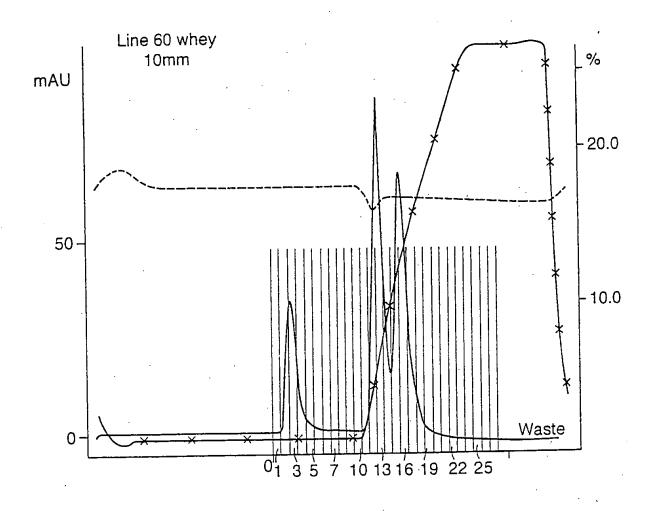
\_\_\_\_\_12099801:1\_UV1\_280nm

\_\_\_\_\_12099801:1\_pH

<del>\_x \_ x \_ x</del> 12099801:1\_Cond%

12099801:1\_Fractions

Fig. 15.

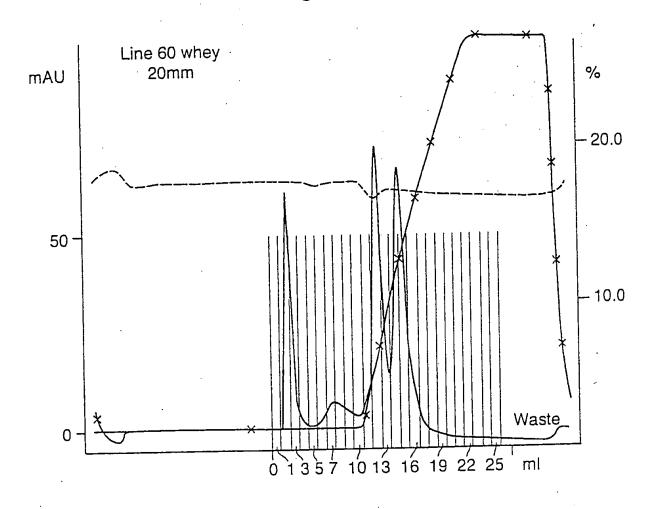


\_\_\_\_\_\_ 12099802:11\_UV1\_280nm

\_\_\_\_\_ 12099802:11\_pH

12099802:11\_Fractions

Fig. 16.



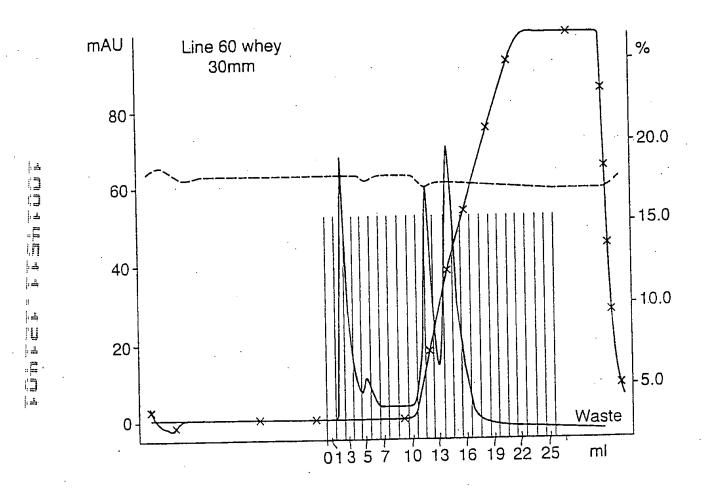
\_\_\_\_\_ 12099803:12\_UV1\_280nm

----- 12099803:12\_pH

<del>-x-x-x-</del> 12099803:12\_Cond%

12099803:12\_Fractions

Fig. 17.



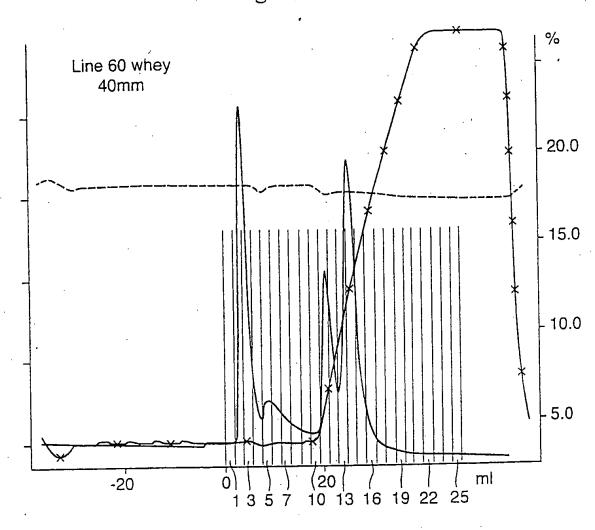
\_\_\_\_\_ 12099804:13\_UV1\_280nm

---- 12099804:13\_pH

<del>× × ×</del> 12099804:13\_Cond%

12099804:13\_Fractions

Fig. 18.



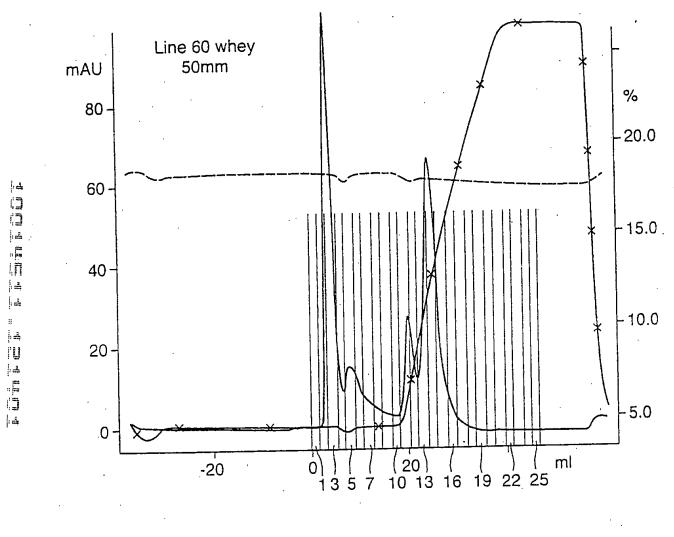
\_\_\_\_\_ 121099805:1\_UV1\_280nm

----- 121099805:1\_pH

<del>-x × × 1</del>21099805:1\_Cond%

121099805:1\_Fractions

Fig. 19.



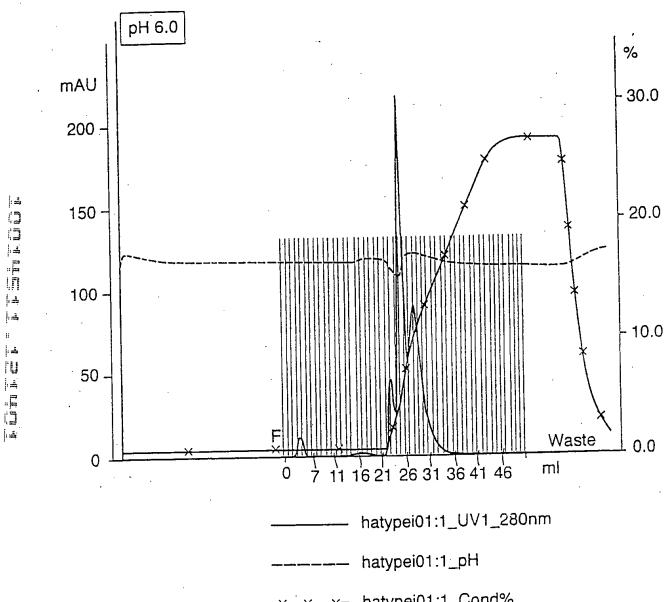
\_\_\_\_\_\_ 121099806:1\_UV1\_280nm

---- 121099806:1\_pH

<del>× × ×</del> 121099806:1\_Cond%

121099806:1\_Fractions

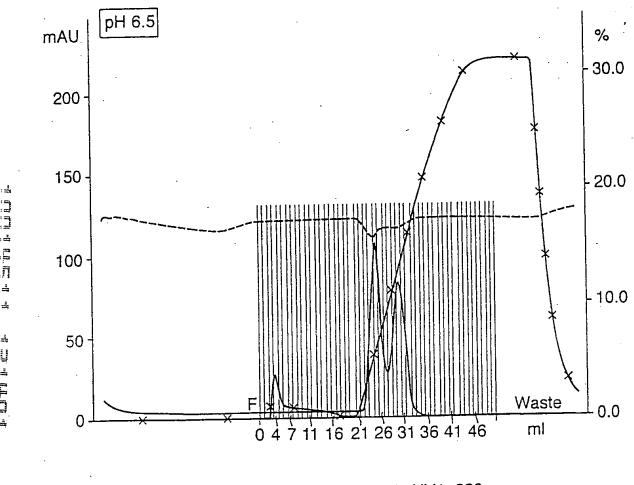
Fig. 20.



hatypei01:1\_Cond%

hatypei01:1\_Fractions

Fig. 21.



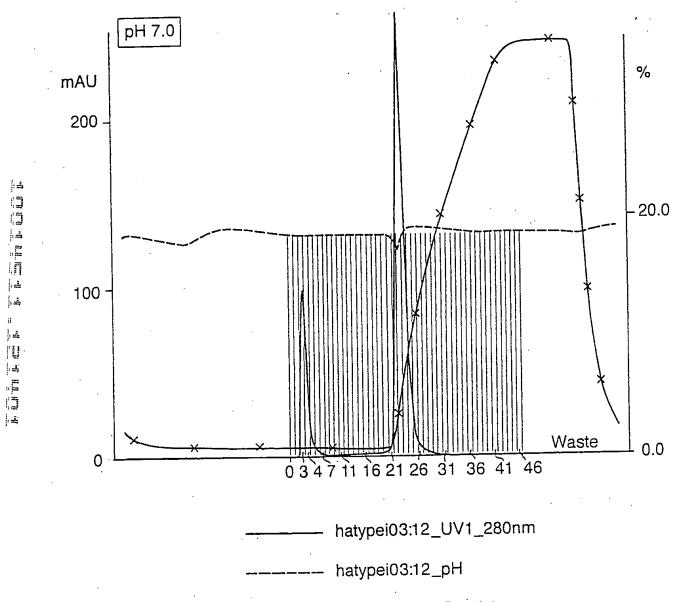
\_\_\_\_\_ hatypei02:11 \_UV1\_280nm

\_\_\_\_\_ hatypei02:11\_pH

-x-x-- hatypei02:11\_Cond%

hatypei02:11 \_ Fractions

Fig. 22.



-x -x -x hatypei03:12\_Cond%

hatypei03:12\_Fractions

Fig. 23.

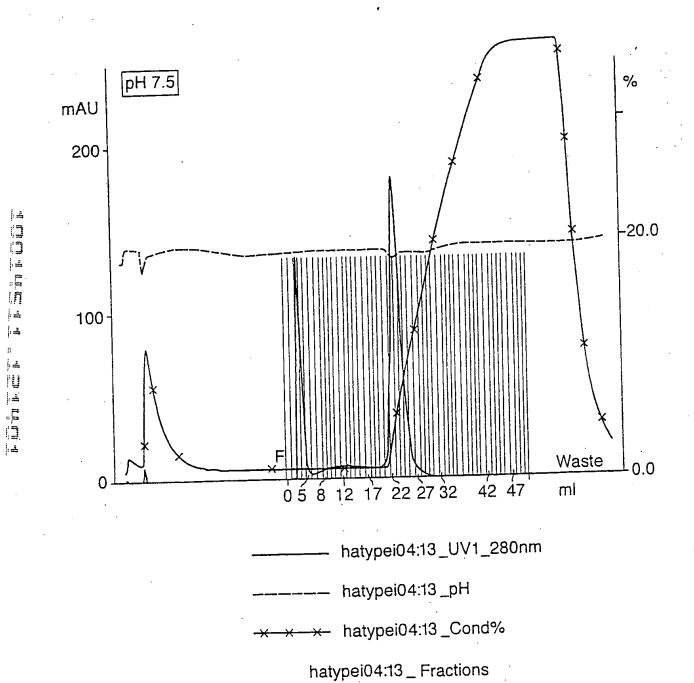
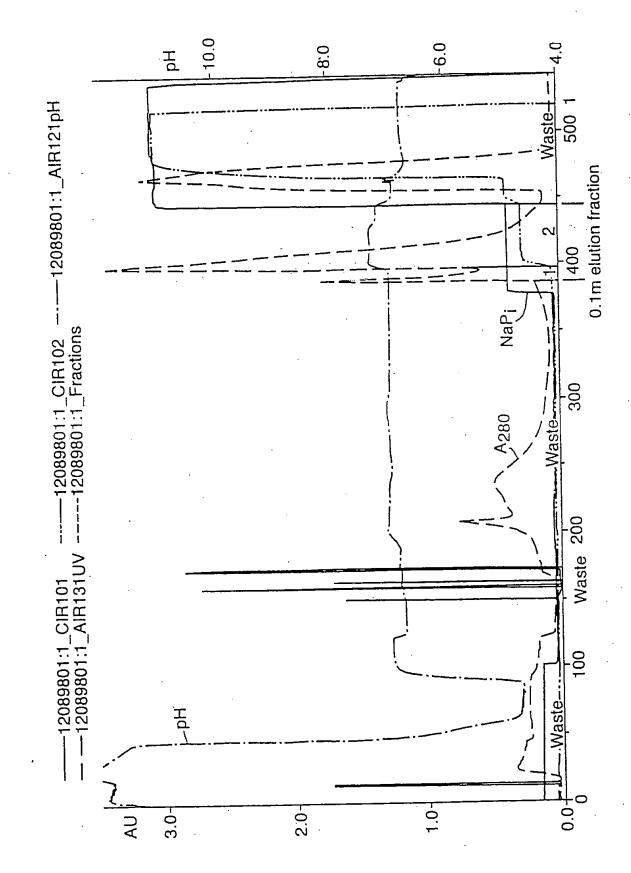
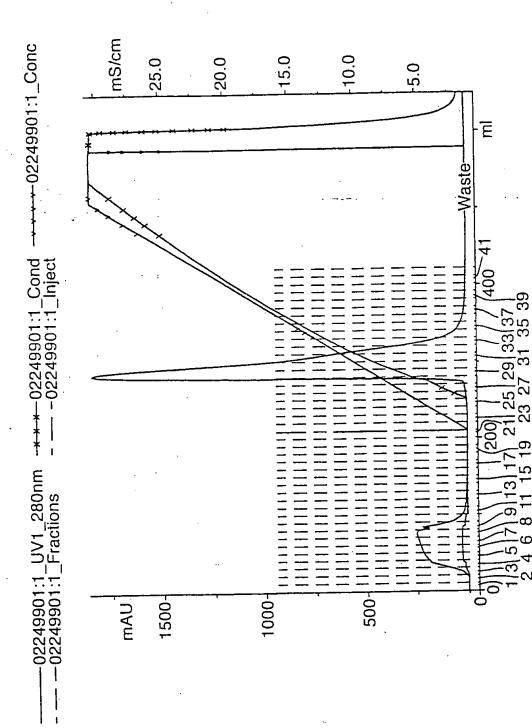


Fig. 24.





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8 11 1

Fig. 26.

XK16/15 80°C cHT type I 10mM Napi pH 6.5 ; QFF eluate Run 02249901/02259901/02269901

1. fr.2-4 2. fr.5-8 3. fr.9-11 1066570 5. fr.2-4 5. fr.2-8 6. fr.5-8 7. fr.9-11 1066570 6. fr.3-8 6. fr.3-Flowthrough

fractions

Sample loaded on CHT